

## p185<sup>neu</sup>-ENCODING DNA AND THERAPEUTICAL USES THEREOF

The present invention relates to plasmid vectors containing p185<sup>neu</sup>-encoding sequences and the use thereof in DNA vaccination against tumours. The plasmids according to the invention contain sequences encoding different fragments of human or rat oncoprotein p185<sup>neu</sup> and are able to induce a 5 humoral or cell-mediated immune response against tumours expressing oncogenes of the ErbB family.

The invention also relates to pharmaceutical compositions containing said plasmids and their use for the prevention or therapy of p185<sup>neu</sup>-expressing tumours.

### 10 **Background of the invention**

Protein p185<sup>neu</sup>, one of the most studied tumour antigens, has raised great interest as target for immune therapy against cancer, due to its presence on the cell membrane of some of the most common human carcinomas.

15 p185<sup>neu</sup> is a membrane receptor encoded in the rat by proto-oncogene Her-2/neu and belonging to the family of Class I Tyrosine Kinase Receptors (RTKs), which also comprises the Epidermal Growth Factor Receptor EGF-R (ErbB-1) and other receptors related thereto (ErbB-3, ErbB-4). These receptors are involved in cell proliferation and differentiation (Hynes and Stern, 1994 BBA 1198:165) and therefore attract a great biological and 20 clinical interest. The receptor consists of three well distinguished domains: an extracellular, transmembrane and intracytoplasmic domain. p185<sup>neu</sup> is involved in the complex network of mechanisms of intracellular signal transduction and intracellular communication that regulate proliferation and differentiation processes (Boyle 1992 Curr. Op. Oncol. 4:156). Oncogene *neu* 25 is named after the chemically-induced rat neuroglioblastoma from which it was first isolated. This activated *neu* form has a single point mutation that

results in the replacement of "A" with "T" and in the consequent substitution of the Valine residue at position 664 of p185<sup>neu</sup> with a glutamic acid residue (Val664Glu) (Bargmann et al. 1986, Cell 45:649).

Also the human *neu* homologous, ErbB-2, has been isolated and characterised and it has been demonstrated that both rat HER2/neu receptor and human ErbB2 have a significant homology with EGFR (Coussens et al. 1985, Sciente 230:1132; Yamamoto et al. 1986, Nature 319:230). While a genetic mutation in the rat sequence is the cause of constitutive receptor activation through dimerization, in ErbB-2 positive human tumours an aberrant expression of the oncogene is observed (Di Marco et al. 1990, Mol. Cell. Biol. 10: 3247; Klapper et al., 2000, Adv Cancer Res, 77:25), even though, in rare cases, activating point mutations and abnormal splicing mechanisms have been found (Kwong et al., 1998, Mol Carcinog, 23:62; Xie et al., 2000, J Natl Cancer Inst, 92:412). The overall effect is similar: gene amplification and increase in the transcription level determine an excess of p185<sup>neu</sup> membrane receptor, with consequent increase of active dimers intracellularly transducing growth signals in a ligand-independent manner. The crystal structure of human and rat p185<sup>neu</sup> extracellular region recently reported shows that this protein is characterised by a rigid conformation that allows to interact with other ErbB receptors, without directly binding any ligands, and trigger the proliferation signal transduction (Cho HS et al. 2003, Nature 421:756).

Under normal circumstances, human p185<sup>neu</sup> is involved in organogenesis and epithelial growth; it is expressed at high levels during placenta formation and fetal development, whereas it is present at very low levels in adult tissues (Press et al. 1990, Oncogene 5:953).

Several studies have demonstrated that human p185<sup>neu</sup> overexpression is associated to the neoplastic process and to the level of tumor aggression. The

overexpression of p185<sup>neu</sup> has been described in lung (Kern et al. 1986, Cancer Res. 50:5184), colon (Cohen et al. 1989, Oncogene 4:81), ovary (Slamon et al. 1989, Science 244:707) adenocarcinomas and in a high number of human mammary carcinomas (Slamon et al. 1989, Science 244:707; Jardines et al. 5 1993, Pathobiology 61:268).

The fundamental properties that make p185<sup>neu</sup> an optimal target for plasmid vaccination are: a) its direct involvement in cell growth and carcinogenesis, therefore clone variants that, due to tumour genetic instability, lose the expression of this antigen also lose their tumorigenicity; b) its 10 expression on the plasmatic membrane, which makes it recognizable by antibodies even in tumour cells that lose the expression of the major histocompatibility system (Lollini P. and Forni G. 2003, Trends Immunol. 24: 62).

Studies carried out on mice transgenic for the activated rat oncogene 15 Her-2/neu (which spontaneously develop p185<sup>neu</sup> positive mammary tumours) and on murine models using p185<sup>neu</sup> positive transplantable tumour lines, have demonstrated the possibility to prevent and cure preneoplastic lesions. As regards in particular the prevention of mammary tumours in mice transgenic for rat activated Her-2/neu, we have demonstrated that the plasmid coding for 20 the extracellular and transmembrane domains of rat p185<sup>neu</sup> is able to induce an in vivo protection more effective than the plasmid encoding for the full-length rat p185<sup>neu</sup> or for the extracellular domain only (secreted antigen) (Amici A. et al. 2000, Gene Ther., 7: 703; Rovero S. et al. 2000, J. of Immunol., 165: 5133). Similar results have been reported by Chen et al. 25 (1998, Cancer Res 58:1965). Other authors have demonstrated that plasmids encoding for p185<sup>neu</sup> - either unvaried or mutated so as to eliminate its tyrosine-kinase activity - are effective in preventing the onset of tumours following to p185<sup>neu</sup>-positive cells inoculum (Wei WZ et al. 1999, Int.

J. Cancer 81: 748). Moreover, plasmids devoid of the signal responsible for the processing through the endoplasmic reticulum (leader), which determines cytoplasmic localization of p185<sup>neu</sup> antigen, proved equally effective. The protection induced by different plasmids was mainly mediated by a humoral immune response in the case of membrane expression of p185<sup>neu</sup>, and by a T-lymphocyte-mediated immune-response in the case of cytoplasmic localization (Pilon SA et al. 2001, J. of Immunol. 167: 3201). However, combined vaccination with plasmids inducing p185<sup>neu</sup> overexpression in both the cytoplasm and the membrane was more effective in protecting against tumour growth (Piechocki MP et al. 2001, J. Immunol. 167: 3367).

Thus, the balance between different immune response mechanisms might be particularly important (Reilly et al., 2001, Cancer Res. 61: 880). Moreover, it has been observed that the vaccination with plasmids encoding for extracellular and transmembrane domains of rat p185<sup>neu</sup> is able to eradicate tumour masses with 2 mm diameter, upon inoculum of cells overexpressing p185<sup>neu</sup>, through a number of different effector mechanisms of the immune system (T helper and T killer cells, antibodies, macrophages, neutrophiles, natural killer cells, Fc receptors, gamma interferon and perforins), which cooperate to tumor rejection (Curcio C. et al. 2003, J. Clin. Invest. 111: 1161).

## 20 Description of the invention

Various constructs encoding the human or human/rat chimeric p185 protein have been inserted in plasmid vectors and used in immunization experiments aimed at preventing tumour progression. For plasmid construction, fragments of the human p185<sup>neu</sup> protein containing the transmembrane domain and portions of the extracellular domain of decreasing length have been prepared from ErbB2 oncogene sequence, or portions thereof have been replaced with homologous sequences from the rat Her-2/neu cDNA so as to create chimeric plasmids.

The plasmids thereby obtained have been evaluated in vaccination experiments in mice inoculated with tumour cells overexpressing human p185<sup>neu</sup>. Plasmids containing truncated forms of p185<sup>neu</sup> induced an antitumor reactivity mediated by killer and helper T lymphocytes, while chimeric 5 plasmids induced an antibody response against both human and rat p185<sup>neu</sup>.

Based on the results of in vivo experiments, the plasmids containing p185<sup>neu</sup> sequences able to induce a strong immune response of both cellular and humoral type have been selected. These plasmids, object of the present invention, contain a sequence encoding a p185<sup>neu</sup> fragment selected from the 10 group consisting of SEQ ID N. 1-14 (human and rat p185<sup>neu</sup> reference sequences are available at Gene Bank accession numbers M11730 and X03362, respectively).

According to the invention, p185<sup>neu</sup> encoding sequences can be inserted in any plasmid vectors suitable for human administration. Besides the 15 encoding sequences, the plasmids can contain functional elements for transcription control, in particular a promoter placed upstream of the encoding sequence, preferably the CMV promoter, start and stop transcription elements, selection markers, such as ampicillin or kanamycin resistance genes, CpG motifs, a polyadenylation site or transcription activators. Transcription control 20 elements should be compatible with the use of the vector in humans. In a preferred embodiment, the plasmids of the invention contain at least 4 CpG motifs, preferably at least 8, up to a maximum of 80. The CpG motifs (ATAATCGACGTTCAA) of bacterial origin induce macrophages to secret IL-12, which in turn induce IFN gamma secretion by natural killer cells, thus 25 activating a T helper lymphocyte-mediated response (Chu R.S. et al. 1997, J. Exp. Med., 186: 1623). Therefore, the insertion of CpG motifs in plasmid sequences enhances the immune response.

In a further embodiment, the invention provides a pharmaceutical

composition containing one or more different plasmids as defined above in association with pharmaceutically acceptable vehicles and excipients. The pharmaceutical compositions, in a form suitable for parenteral administration, preferably in the form of injectable solution, are conveniently used for DNA 5 vaccination. Principles and methods for DNA vaccination are known to the skilled in the art and are disclosed, for example, in Liu MA 2003; J Int Med 253: 402.

In another embodiment, the invention provides a combined preparation containing at least two, preferably at least four, more preferably at least eight 10 different plasmids for simultaneous, sequential or separate administration to a subject or patient.

Plasmids, compositions and preparations according to the invention are used in preventive or therapeutical treatment of subjects at risk of developing p185<sup>neu</sup>-positive tumours, or patients with primary tumours, metastasis or 15 relapses of p185<sup>neu</sup>-positive tumours. Prevention can be primary, when the tumour is not manifest, secondary, when the tumour is in the initial phases as a preneoplastic lesion, or tertiary, in the case of tumour relapse or metastatic process. Tumours that can benefit from treatment with the plasmids of the 20 invention are those of epithelial origin, in particular pulmonary, ovary and mammary adenocarcinomas and, more generally, tumours expressing the p185<sup>neu</sup> protein.

#### **Detailed description of the invention**

##### *Construction of the plasmid backbone of pCMV3.1*

To construct plasmids encoding human p185<sup>neu</sup> fragments and chimeric 25 plasmids, the pCMV3.1 plasmidic backbone was used. Fragments deriving from human proto-oncogene ErbB-2 cDNA and from rat proto-oncogene Her-2/neu cDNA have been inserted in pCMV3.1 (Invitrogen, Milano, Italia) by removing with restriction enzymes DraIII (nt1531) e BsmI (nt3189) a

fragment of 1658 bp containing the replication origin f1, the replication origin and the early SV40 promoter, the gene encoding for neomycin resistance and SV40 polyadenylation signal. The resulting modified plasmid (pCMV3.1) present some advantages compared to native pcDNA3.1. In fact, the size 5 reduction to 3900 bp and the removal of irrelevant sequences contribute to increase transfection efficacy *in vivo*.

*Construction of plasmid pCMV3.1erbB2*

Human ErbB2 cDNA, obtained from plasmid pSVerbB2, has been inserted in the multiple cloning site of pCMV3.1 at restriction sites HindIII 10 and XbaI. This plasmid is used for the construction of plasmids expressing truncated p185<sup>neu</sup> and chimeric plasmids.

*Construction of plasmids containing the sequence 4XCpG: pCMV3.1hECD-TM-4CpG and pCMV3.1hECD-TM-4noCpG*

After removal of the sequence encoding the intracytoplasmic domain 15 from plasmid pCMV3.1-erbB2, two plasmids coding for proto-oncogene ErbB2 extracellular and transmembrane regions were prepared. The procedure comprised first the restriction analysis to identify the unique sites present in the nucleotide sequence of ErbB2 cDNA. A unique site recognized by enzyme AccIII (nt 2195) about 20 bp downstream of the end of the transmembrane 20 domain was identified.

The cytoplasmic domain was removed using the enzyme AccIII present as unique restriction site and enzyme XbaI. To re-insert at the 3'end of the DNA of the ErbB2 ECD-TM the nucleotide triplet TAA, recognized as translation stop signal, we used two synthetic sequences consisting of two sense 25 (oligonucleotide #1, #3) and antisense (oligonucleotide #2, #4) oligonucleotides having the restrictions sites AccIII and XbaI at their ends. In these synthetic sequences there are also four repeated sequences CpG and noCpG. The latter is used as negative control. These two new plasmids have

been named pCMV3.1hECD-TM-4CpG and pCMV3.1hECD-TM-4noCpG.

*Construction of the plasmids containing the sequence 8XCpG: pCMV3.1H/NhECD-TM-8CpG and pCMV3.1H/NhECD-TM-8noCpG*

To add further aspecific immune stimuli we constructed a new plasmid  
5 backbone containing 4 immune-stimulating CpC sequences, called pCMV3.1  
H/N-4CpG. For this purpose we modified pCMV3.1 so as to remove one of  
the two restriction sites for the enzyme PmeI and invert the restriction sites for  
HindIII and NneI present on the multiple cloning site by means of a synthetic  
sequence consisting of two sense (oligonucleotide #5) and antisense  
10 (oligonucleotide #6) oligonucleotides. In this new plasmid, named pCMV3.1  
H/N, two synthetic sequences have been inserted, consisting of two sense  
(oligonucleotide #7, #9) and antisense nucleotides (oligonucleotide #8, #10),  
containing four repeats for the CpG and noCpG sequences in the unique  
restriction sites XbaI and PmeI, thus obtaining pCMV3.1 H/N-4CpG and  
15 4noCpG. Thereafter, DNA fragments hECD-TM-4CpG and hECD-TM-  
4noCpG have been inserted in pCMV3.1 H/N-4CpG and in pCMV3.1  
H/N-4noCpG respectively, thus obtaining two new plasmids called  
pCMV3.1H/N-hECD-TM-8CpG and pCMV3.1H/N-hECD-TM-8noCpG.

*Construction of the plasmid containing the sequence of the second  
20 cystein domain and transmembrane domain of human p185<sup>neu</sup>:  
pCMV3.1H/Nh2°cysECD-TM-8CpG*

Human p185<sup>neu</sup> extracellular domain is characterised by two regions  
rich in cysteins, known as 1<sup>st</sup> and 2<sup>nd</sup> cystein sub-domain (1<sup>st</sup> cys and 2<sup>nd</sup> cys).  
Unlike the rat cDNA sequence containing only one site BstEII (nt1250) in the  
25 extracellular domain, located in the nucleotide region that separates 1<sup>st</sup> cys  
from 2<sup>nd</sup> cys, the cDNA sequence of the extracellular domain of ErbB2 has  
two restriction sites for BstEII: in addition to the site in the same position as  
that of rat (nt1372), a further BstEII site (nt963) is present in the portion

encoding the 1<sup>st</sup> cys of the extracellular domain. Digesting plasmid pCMV3.1H/NhECD-TM-8CpG with HindIII and BstEII, a DNA fragment consisting of the 2nd cys from the extracellular domain, the transmembrane domain, the sequence 8CpG and the plasmid pCMV3.1H/N was obtained.

5 Then the signal for rat p185neu secretion through the endoplasmic reticulum was inserted by enzymatic DNA amplification (PCR reaction) using a sense oligonucleotide consisting of the primer T7 (oligonucleotide #11) which recognizes the T7 RNA polymerase, present at the beginning of the pCMV3.1H/N multiple cloning site, and an antisense oligonucleotide 10 (oligonucleotide #12) having the BstEII site at its end. After purification, enzymatic digestion of the amplified fragment with restriction enzymes HindIII and BstEII and subsequent cloning, pCMV3.1H/Nh2°cys-TM-8CpG (Fig. 1) has been obtained. (Fig. 1). This plasmid was used in vaccination experiments, to have it compared with pCMV3.1 H/NhECD-TM-8CpG. 15 Thereafter, a chimeric cDNA encoding for the fusion protein between 2<sup>nd</sup> cys and transmembrane domain (nt 1372-nt 2204) of the human sequence and 1<sup>st</sup> cys (nt 1-nt 1250) of the rat sequence has been prepared. The reconstitution of the entire protein sequence by the fusion of portions deriving from rat and human cDNAs, respectively, allows to increase the immune response.

20 *Construction of the chimeric plasmid containing the sequence of the first cystein domain of rat p185neu and of the second cystein domain and transmembrane domain of human (nt 1-nt 1250): pCMV3.1H/N-r1°cys-h2°cysTM-8CpG*

Unlike the rat cDNA sequence containing only a BstEII (nt1250) site in 25 the extracellular domain located in the nucleotide region that separates the first and the second region rich in cysteins, the cDNA sequence of the extracellular domain of Erb2 has two restriction sites for BstEII: one in position 1372 (nt), as in the rat sequence, and the other in position 963 (nt),

i.e. in the sequence portion encoding for the 1<sup>st</sup> cys of the extracellular domain. The presence of the BstEII site in the same position both in the rat cDNA domain (1250nt) and in the human cDNA (1372nt) allowed the construction of a plasmid able to encode a fusion product between rat 1<sup>st</sup> cys and human 2<sup>nd</sup> cys. In fact, digesting pCMV3.1H/N-h2°cysTM-8CpG with restriction enzymes HindIII and BstEII allowed to replace the DNA fragment encoding for rat p185<sup>neu</sup> secretion signal with the nucleotide sequence encoding for rat 1st cys obtained through digestion of pCMV3.1rECD-TM-4CpG with the same enzymes. The product of plasmid pCMV3.1H/N-r1°cys-h2°cysTM-8CpG (Fig. 2) consists of a portion of 412 aa of rat p185neu and a portion of 274 aa of human p185<sup>neu</sup>. This new plasmid, pCMV3.1H/Nr1°cys-h2°cysTM-8CpG has been used in vaccination experiments using pCMV3.1H/N-hECD-TM-8CpG as comparative term. Surprisingly, the plasmid coding for the chimeric protein induces in mice a complete protection against tumours expressing human p185<sup>neu</sup> (Table). This protection is similar to that induced by pCMV3.1H/N-hECD-TM-8CpG. Moreover, analysis of the sera of mice vaccinated with both plasmids has evidenced a similar antibody titer towards human p185<sup>neu</sup>.

*Plasmids able to encode decreasing fragments of the extracellular and transmembrane domain of human p185 neu*

Construction of seven plasmids that encode decreasing fragments of the extracellular and transmembrane domain of human p185<sup>neu</sup>, namely: pCMV3.1H/NhECD1-TM-8CpG (-70 aa), pCMV3.1H/NhECD2-TM-8CpG (-150 aa), pCMV3.1H/NhECD3-TM-8CpG (-230 aa), pCMV3.1H/NhECD4-TM-8CpG (-310 aa), pCMV3.1H/NhECD5-TM-8CpG (-390 aa), pCMV3.1H/NhECD6-TM-8CpG (-470 aa) and pCMV3.1H/NhECD7-TM-8CpG (-550 aa).

The fragment encoded by the first of these fragments is 70 aa (deletion

of 360 bp) shorter. All the others are gradually 80 aa shorter (deletions of 240 bp).

These fragments have been obtained by DNA enzymatic amplification, using seven different sense oligonucleotides with NheI restriction site (oligonucleotides #13-#19) at its end and an antisense oligonucleotide (oligonucleotide #20) able to recognise the site called "pcDNA3.1/BGH Reverse priming site" (830-850 nt) present at the 3' end of the polylinker of pCMV3.1. Further to enzymatic digestion with restriction enzymes NheI and PmeI, amplification products have been cloned in pCMV3.1H/N-neu leader, 10 previously obtained inserting the secretion signal to the endoplasmic reticulum of rat p185<sup>neu</sup> in restriction sites. The DNA fragment of rat p185<sup>neu</sup> secretion signal has been obtained by enzymatic DNA amplification using primer T7 (oligonucleotide #11) as sense nucleotide and an antisense nucleotide (oligonucleotide #21) with NheI site at its end. The amplified fragment after 15 purification and restriction digestions with HindIII and NheI has been cloned in plasmid pCMV3.1H/N, digested with the same enzymes, thus obtaining the pCMV3.1H/N-neu leader. Membrane expression of the different truncated forms of human p185<sup>neu</sup> is expected in view of the presence of the secretion signal to the endoplasmic reticulum of rat p185<sup>neu</sup>. The plasmids encoding the 20 truncated forms pCMV3.1H/NhECD1-TM-8CpG (Fig. 3), pCMV3.1H/NhECD2-TM-8CpG (Fig. 4), pCMV3.1H/NhECD3-TM-8CpG (Fig. 5), pCMV3.1H/NhECD4-TM-8CpG (Fig. 6) as well as the control plasmid pCMV3.1H/NhECD-TM-8CpG, protect 100% of the vaccinated mice against a lethal inoculum of tumour cells expressing human p185<sup>neu</sup> (Table). 25 Plasmid pCMV3.1H/NhECD5-TM-8CpG (Fig. 7) protects 60% of the animals (Table), while plasmids pCMV3.1H/NhECD6-TM-8CpG and pCMV3.1H/NhECD7-TM-8CpG (Fig. 8, 9), do not have protective effect against a lethal inoculum of tumour cells expressing human p185<sup>neu</sup> (Table).

The protein products expressed by the different plasmids are not secreted through the endoplasmic reticulum. The absence of consensus sequences necessary for glycosilation and for their processing through the endoplasmic reticulum, or conformational changes due to deletion of aminoacids at the -

5 NH<sub>2</sub> terminus, could explain the absence of protein products in the membrane. Therefore, to further verify if the various truncated forms of the extracellular and transmembrane domain of human p185<sup>neu</sup> were correctly expressed, new plasmids coding for fusion proteins characterized by epitope myc at the -NH<sub>2</sub> terminus were generated. These recombinant proteins are recognized by an

10 anti-myc monoclonal antibody, therefore it is possible to analyse their expression and localisation by confocal microscopy.

First a new plasmid coding for the secretion signal to the rat endoplasmic reticulum (neu leader) and for the myc epitope has been created. Cloning has been carried out using a synthetic sequence consisting of a sense

15 (oligonucleotide #22) and antisense (oligonucleotide #23) having at both ends the NheI site. The NheI site in the 5' position was mutated so that, once correctly ligated, it was not recognized by the enzyme. We thus obtained the pCMV3.1H/Nneuleader-myc epitope. With this plasmid, the sequences encoding human p185<sup>neu</sup> truncated forms have been cloned in the restriction

20 sites NheI and PmeI. Then, 3T3 NIH fibroblasts have been transfected in vitro with plasmids using lipofectamine 2000 (Invitrogen, Milan, Italy). After 48 hours the transfected cells have been analysed with confocal microscopy, using a FITC-conjugated anti-myc monoclonal antibody (Sigma-Aldrich Srl, Milan, Italy). It has been thus demonstrated that all the pasmids-encoded

25 truncated forms are located in the cytoplasm. 3T3 NIH fibroblasts have been transfected in parallel with plasmid pCMV3.1H/NhECD-TM-8CpG and analysed with confocal microscopy using c-erbB2/c-neu Ab-3 monoclonal antibody (Oncogene, Boston, MA) as primary antibody and a

FITC-conjugated anti-mouse secondary antibody (PharMigen, San Diego, CA). It was thus observed that human ECD-TM is expressed in the membrane. The results obtained using the first four plasmids described previously (pCMV3.1H/NhECD1-TM-8CpG, pCMV3.1H/NhECD2-TM-8CpG, 5 pCMV3.1H/NhECD3-TM-8CpG, pCMV3.1H/NhECD4-TM-8CpG), demonstrate that a cellular response is sufficient for antitumour prevention. However, it is known that contemporaneous activation of the cellular and humoral response is necessary for a more effective therapy (Rielly et al., 2001, Cancer Res 61:880). As already described in the previous paragraph, the 10 chimeric protein encoded by plasmid pCMV3.1H/N-r1°cys-h2°cysTM-8CpG is able to protect 100% of the vaccinated animals and is able to induce a strong humoral response in the mice.

*Chimeric plasmids able to encode for five different man-rat chimeric p185<sup>neu</sup>*

15 For the construction of plasmids coding for chimeric proteins, we selected pCMV3.1H/NhECD1-TM-8CpG, pCMV3.1H/NhECD2-TM-8CpG, pCMV3.1H/NhECD3-TM-8CpG and pCMV3.1H/NhECD4-TM-8CpG. These four plasmids protect 100% of the vaccinated mice against a lethal inoculum of tumour cells expressing human p185<sup>neu</sup>. Also plasmid 20 pCMV3.1H/NhECD5-TM-8CpG has been selected, even if it protects only 60% of the vaccinated mice, because the encoded protein differs only by 17 aa from that encoded by pCMV3.1H/Nh2°cysECD-TM-8CpG (275 aa), which protects 20% of the vaccinated mice. We can hypothesize that the peptide sequence of 17 aa corresponds to an important epitope for the induction of an 25 effective immune response.

DNA fragments encoding for rat p185<sup>neu</sup> portions have been obtained by DNA enzymatic amplification. To amplify these cDNA fragments six oligonucleotides having all the same orientation, namely that of T7 primer

(oligonucleotide #11), have been used, while the five antisense have been designed to recognize rat cDNA in the proper positions and have the restriction site for NheI at their ends (oligonucleotides #24-#28). After purification and digestion with restriction enzymes HindIII and NheI, the 5 amplified fragments have been inserted in the corresponding plasmids (pCMV3.1H/NhECD1-TM-8CpG, pCMV3.1H/NhECD2-TM-8CpG, pCMV3.1H/NhECD3-TM-8CpG, pCMV3.1H/NhECD4-TM-8CpG pCMV3.1H/NhECD5-TM-8CpG) and digested with the same restriction enzymes. In this way we obtained five new plasmids able to code for chimeric 10 proteins of 689 aa, of which 2 (Val-Ser) belong to restriction site NheI used for the conjunction between rat and human DNA. The presence of these two aa renders both human and rat portions heteroclytic.

The chimeric proteins differ for human p185<sup>neu</sup> decreasing portions and rat p185<sup>neu</sup> increasing portions. Plasmid pCMV3.1H/Nr73-hECD1-TM-8CpG 15 (Fig. 10) encodes 73 aa of the rat p185<sup>neu</sup> extracellular domain and 614 aa of human p185<sup>neu</sup>. Plasmid pCMV3.1H/Nr153-hECD2-TM-8CpG (Fig. 11) encodes 153 aa of the rat p185<sup>neu</sup> extracellular domain and 534 aa of human p185<sup>neu</sup>. Plasmid pCMV3.1H/Nr233-hECD3-TM-8CpG (Fig. 12) encodes 233 aa of the rat p185<sup>neu</sup> extracellular domain and 454 aa of human p185<sup>neu</sup>. 20 Plasmid pCMV3.1H/Nr313-hECD4-TM-8CpG (Fig. 13) encodes 313 aa of the rat p185<sup>neu</sup> extracellular domain and 374 aa of human p185<sup>neu</sup>. Plasmid pCMV3.1H/Nr393-hECD5-TM-8CpG (Fig. 14) encodes 393 aa of the rat p185<sup>neu</sup> extracellular domain and 294 aa of human p185<sup>neu</sup>. Indirect evidence 25 of the membrane expression of human/rat chimeric p185<sup>neu</sup> encoded by these plasmids has been obtained immunizing mice with the five new plasmids and with pCMV3.1H/N-r1°cys-h2°cysTM-8CpG as positive control. The sera of all vaccinated mice contain specific antibodies against human p185<sup>neu</sup>. Moreover, the animals vaccinated with plasmids encoding the five different

chimeric proteins are also protected with a lethal inoculum of tumour cells expressing human p185<sup>neu</sup>.

## EXAMPLES

### Example 1 - Construction of plasmid pCMV3.1H/N-r1°cys-h2°cysTM-

#### 5 8CpG

To construct chimeric plasmid pCMV3.1H/N-r1°cys-h2°cysTM-8CpG we started from plasmid pCMV-ECD-TM, which expresses the extracellular and transmembrane domain of rat p185<sup>neu</sup> (Amici et al 2000, Gene Ther., 7: 703). pCMV-ECD-TM was digested with restriction enzymes HindIII and 10 XbaI (BioLabs, Beverly, MA) to separate the insert from the plasmid backbone.

#### *Restriction digestion with enzyme HindIII:*

|                               |                     |
|-------------------------------|---------------------|
| plasmid DNA (1 µg/µl)         | 10 µl               |
| restriction buffer 10X (NEB2) | 10 µl               |
| 15 HindIII (10U/µl)           | 5 µl                |
| H <sub>2</sub> O              | <u>75 µl</u>        |
|                               | 100 µl final volume |

The mixture was incubated at 37°C for 4 hours and the digestion product controlled by electrophoresis on 1% agarose gel using a molecular 20 weight marker and undigested plasmid as control.

Once confirmed plasmid linearization, DNA was precipitated by adding 1/10 volume of 3 M sodium acetate at pH 5.2 and 2 volumes of cold absolute ethanol.

The sample was kept on ice for 20 min., then centrifugated with a 25 minicentrifuge at 14.000 rpm for 12 min. The pellet was washed three times with 1 ml 70% cold ethanol, dried under vacuum for 5 min, then resuspended in 84 µl H<sub>2</sub>O and enzymatically digested with restriction enzyme XbaI.

*Restriction digestion with enzyme XbaI:*

|   |             |
|---|-------------|
| DNA resuspended in H <sub>2</sub> O (10 µg) | 84 µl       |
| Restriction buffer 10X (NEB2)               | 10 µl       |
| BSA 100X (100mg/ml)                         | 1 µl        |
| 5 XbaI (10U/ml)                             | <u>5 µl</u> |
|   | 100 µl      |

The mixture was incubated at 37°C for 4 hours and the digestion product was precipitated and dried as described above. DNA was resuspended in 30 µl H<sub>2</sub>O.

10 The two DNA fragments corresponding to the plasmid backbone (pCMV of 4400bp) and to the insert (ECD-TM of 2100bp) were separated by electrophoresis on a 1% agarose gel.

The band corresponding to the insert was removed and DNA eluted from the gel using a Qiaquick gel extraction kit (Qiagen Italy).

15 In parallel, the new plasmid backbone (pCMV3.1H/N-4CpG) wherein the DNA fragment corresponding to rat p185 ECD-TM, was digested with the same restriction enzymes and eluted on agarose gel.

20 The DNA fragments corresponding to rat ECD-TM and the linearized plasmid pCMV3.1H/N-4CpG were used to obtain pCMV3.1H/N-rECD-TM-4CpG by ligation reaction.

*Ligation reaction*

|  |             |
|--|-------------|
| DNA insert (rECD-TM) (50 ng/µl)                    | 2 µl        |
| Linearized plasmid DNA (pCMV3.1H/N4CpG) (50 ng/µl) | 1 µl        |
| Reaction buffer 10X for T4 DNA ligase              | 1 µl        |
| 25 T4 DNA ligase (2U/µl)                           | 1 µl        |
| H <sub>2</sub> O                                   | <u>5 µl</u> |
|  | 10 µl       |

The ligation reaction was incubated at 16°C for 4 hours.

The ligation product was then used to transform the *E. coli* bacterial strain DH5  $\alpha$ . The bacterial cells have been made competent with the CaCl<sub>2</sub> technique.

Transformation of the bacterial strain DH5  $\alpha$ :

|   |                           |             |
|---|---------------------------|-------------|
| 5 | Competent bacterial cells | 100 $\mu$ l |
|   | Ligation product          | 5 $\mu$ l   |

To make the plasmid DNA penetrate the competent cells, these were kept on ice for 40 min. and submitted to thermal shock (1.5 min. at 42°C and then 2 min. on ice).

10 After adding 1 ml LB growth medium, the transformed bacterial cells were incubated at 37°C for 1 hour to restore their physiological conditions.

The cell suspension was then centrifuged at 6000 rpm for 1 min. and the pellet was resuspended in 100  $\mu$ l LB.

15 The cells were seeded in Petri dishes containing selective solid medium (LB with agar + ampicillin 100  $\mu$ g/ml) and grown at 37°C for 1 night. Ampicillin allows the growth of cells containing plasmid pCMV3.1H/N-rECD-TM-4CpG which confers ampicillin-resistance.

The resulting clones were analysed by alkaline lysis to select those containing the recombinant plasmid pCMV3.1H/N-rECD-TM-4CpG.

20 To obtain chimeric plasmid pCMV3.1H/N-r1°cys-h2°cysTM-8CpG, plasmid pCMV3.1H/N-rECD-TM-4CpG was digested with restriction enzymes BstEII and XbaI to remove the second cystein domain together with the transmembrane domain of rat p185<sup>neu</sup>. At the same time, plamid pCMV3.1hECD-TM-4CpG was digested with the same enzymes to isolate the 25 DNA fragment corresponding to the second cystein subdomain and transmembrane domain of the human gene.

*Digestion with BstEII:*

|                               |                     |
|-------------------------------|---------------------|
| plasmid DNA (1 µg/µl)         | 10 µl               |
| Restriction buffer 10X (NEB3) | 10 µl               |
| BstEII (10U/µl)               | 5 µl                |
| 5 H <sub>2</sub> O            | <u>75 µl</u>        |
|                               | 100 µl final volume |

The mixture was incubated at 60°C for 4 hours.

Restriction digestion with XbaI, recovery of the fragments to be used for cloning, ligation reaction and transformation of competent cells have been described previously.

The resulting chimeric plasmid pCMV3.1H/N-r1°cys-h2°cysTM-8CpG has been sequenced using the automatic ABI PRISM 310 Genetic Analyzer (Applied Biosystem), to verify the correct insertion of the fragment corresponding to the 2<sup>nd</sup> cystein subdomain and the transmembrane domain of the human gene.

Example 2 - in vivo test*Animals*

Balb/cAnCr (H-2<sup>d</sup>) female mice aged about seven weeks have been used for all experiments.

20 The animals, supplied by Charles River Laboratories (Calco, MI, Italy), are grown in aseptic conditions and in accordance with the European Community guidelines.

*Intramuscular administration followed by in vivo electroporation*

To avoid unwanted contractions of the tibial muscle each mouse was 25 anaesthetized by i.p. inoculum of 300 µl avertine, made of 0.58 g tribromoethanol (Sigma-Aldrich) and 310 µl Tert-Amyl alcohol (Aldrich) dissolved in 39.5 ml deionized H<sub>2</sub>O. All mice have been then shaved in correspondence of the tibial muscle for the inoculum.

The animals have been vaccinated in correspondence of both antero-tibial muscles, with 40 µl of solution containing 50 µg DNA.

The DNA-containing mixture was prepared shortly before use, in conformity with the indications of Dr. F. Pericle (Valantis, Inc., The 5 Woodlands, Texas, USA). This solution contains 1.25 mg/ml plasmid DNA, 6 mg/ml poly-L-glutamate sodium salt (Sigma-Aldrich, S.r.l., Milano, Italia), 150 mM sodium chloride (Fluka, BioChemika, Buchs, Switzerland) and distilled water free from endotoxins (Nucleare Free Water, Promega Corporation) to a final volume of 1 ml.

10 After about 5 min from the inoculum, the treated area was submitted to electroporation, by application of two electric impulses having an intensity of 375 V/cm<sup>2</sup>, each lasting 25 ms, using the electroporator Electro Square Porator (T820, BTX, San Diego, CA, USA). The transcutaneous electric impulses have been applied by use of two square steel electrodes placed at 15 3 mm from each other, beside each paw. Gene immunization by electroporation was carried out twice for each animal 21 and 7 days before inoculum of tumour cells.

*Inoculum of tumour cells*

20 The mice have been inoculated with a suspension containing 2 x 10<sup>5</sup> D2F2/E2 cells. These cells derive from a mammary tumour spontaneously generated in a hyperplastic alveolar node of a BALB/c mouse and express high levels of human p185.

*In vivo evaluation of tumour growth*

25 Tumour growth was evaluated weekly by palpation and the dimensions of the tumours were measured along two perpendicular diameters with a calibre. Neoplastic masses measuring more than 3 mm are considered as tumours.

Tumour growth was followed for 100 days from tumour inoculum or

until the tumour had grown to a diameter higher than 10 mm, then animals were sacrificed.

**Table**

Mice: female BALB/c

5 Tumour: D2F2-E2 expressing human p185<sup>neu</sup>

| plasmids                          | Number of mice | protection | antibodies |
|-----------------------------------|----------------|------------|------------|
| pCMV3.1H/N-8CpG                   | 5              | 0%         | -          |
| pCMV3.1H/N-hECD-TM-8CpG           | 5              | 100%       | +++        |
| pCMV3.1H/N-hECD1-TM-8CpG          | 5              | 100%       | -          |
| pCMV3.1H/N- hECD2-TM-8CpG         | 5              | 100%       | -          |
| pCMV3.1H/N- hECD3-TM-8CpG         | 5              | 100%       | +          |
| pCMV3.1H/N- hECD4-TM-8CpG         | 5              | 100%       | ++         |
| pCMV3.1H/N- hECD5-TM-8CpG         | 5              | 60%        | -          |
| pCMV3.1H/N- hECD6-TM-8CpG         | 5              | 0%         | -          |
| pCMV3.1H/N- hECD7-TM-8CpG         | 5              | 0%         | -          |
| pCMV3.1H/N-r1°cys-h2°cys.-TM-8CpG | 5              | 100%       | +++        |

**List of oligonucleotides synthesized and used for plasmid construction**

#1. AccIII-TAA-4CpG-erbB2 sense 71 nt

10 5'CCGGAAGTAAATAATCGACGTTCAAATAATCGACGTTCAAAT  
AATCGACGTTCAAATAATCGACGTTCAAT 3'

#2. XbaI-TAA-4CpG-erbB2 antisense 71 nt

5'CTAGATTGAACGTCGATTATTGAACGTCGATTATTGAACG  
TCGATTATTGAACGTCGATTATTACTT 3'

#3. AccIII-TAA-4noCpG-erbB2 sense 71 nt

5'CCGGAAGTAAATAATAGAGCTCAAATAATAGAGCTC~~AAA~~

5 TAATAGAGCTCAAATAATAGAGCTCAAT 3'

#4. XbaI-TAA-4noCpG-erbB2 antisense 71 nt

5'CTAGATTGAAGCTCTATTATTGAAGCTCTATTATTGAA~~GCT~~  
CTATTATTGAAGCTCTATTATTACTT 3'

#5. HindIII-NheI sense 27nt

10 5' CTAGGAAGCTTGTAACTTGCTAGCT 3'

#6. HindIII-NheI antisense 27 nt

5'AGCTAGCTAGCAAGTTAACAAAGCTTC 3'

#7. XbaI-4CpG-neu sense 68 nt

5'CTAGATAATCGACGTTCAAATAATCGACGTTCAAATAATCGA  
15 CGTTCAAATAATCGACGTTCAAGTTT 3'

#8. PmeI-CpG-neu antisense 64 nt

5'AAACTTGAACGTCGATTATTGAACGTCGATTATTGAAC GT  
CGATTATTGAACGTCGATTAT 3'

#9. XbaI-4noCpG-neu sense 68 nt

20 5'CTAGATAATAGAGCTCAAATAATAGAGCTCAAATAATA~~AG~~  
AGCTTCAAATAATAGAGCTTCAAGTTT 3'

#10. PmeI-4noCpG-neu antisense 64 nt

5'AAACTTGAAGCTCTATTATTGAAGCTCTATTATTGAAGCT  
CTATTATTGAAGCTCTATTAT 3'

25 #11. T7 primer

5'TAATACGACTCACTATAGGG 3'

#12. BstEII-neuleader antisense 32 nt

5'GGCCGGTTACCCGCGATTCCGGGGGCAGGAG 3'

#13. hECD1-TM-sense-NheI 35 nt

5'CCGGCTAGCTAGCCTGTCCTCCTGCAGGATATCC 3'

#14. hECD2-TM-sense-NheI 35 nt

5'CCGGCTAGCTAGCGGAGGGTCTGATCCAGCGGA 3'

5 #15. hECD3-TM-sense-NheI 35 nt

5'CCGGCTAGCTAGCCTGCCACTGACTGCTGCCATG 3'

#16. hECD4-TM-sense-NheI 35 nt

5'CCGGCTAGCTAGCTGCACCCCTCGTCTGCCCTGC 3'

#17. hECD5-TM-sense-NheI 35 nt

10 5'CCGGCTAGCTAGCCCGCTCCAGCCAGAGCAGCTCC 3'

#18. hECD6-TM-sense-NheI 35 nt

5'CCGGCTAGCTAGCAACACCCACCTCTGCTTCGTGC 3'

#19. hECD7-TM-sense-NheI 35 nt

CCGGCTAGCTAGCCCCAGGGAGTATGTGAATGCCA 3'

15 #20. pcDNA3.1/BGH Reverse primer 20 nt

5'TAGAAGGCACAGTCGAGGCT 3'

#21. NheI-neuleader-antisense 43 nt

5'CCGGCTAGCTAGCCCGATTCCGGGGGCAGGAGGGCGAGG  
AG 3'

20 #22. His-myc-sense-noNheI 69 nt

5'CTAGGCATCATCATCATCATCATATAATGGTCATACCGGTGAAC  
AAAAACTCATCTCAGAAGAGGGATCTGG 3'

#23. His-myc-antisense-NheI 69 nt

5'CTAGCCAGATCCTCTTGAGATGAGTTTTGTTCACCGGTAT  
25 GACCATTATGATGATGATGATGC 3'

#24. NheI-73neu antisense 35 nt

5'CCGGCTAGCTAGCGCTGGCATTGGCAGGCACGTAG 3'

#25. NheI-153neu antisense 35 nt

5'CCGGCTAGCTAGCCAGGATCTCTGTGAGACTTCGA 3'

#26. NheI-233neu antisense35 nt

5'CCGGCTAGCTAGCGCCCTTGCACCGGGCACAAACCA 3'

#27. NheI-313neu antisense35 nt

5'CCGGCTAGCTAGCTCCACTTCCGTAGACAGGTAG 3'

#28. NheI-393neu antisense 35 nt

5'CCGGCTAGCTAGCAATGCCGGAGGAGGGGTCCCCA 3'